AMENDMENT

Please amend the application as follows:

In the specification:

Figure 1. A) DNA (SEQ ID NO:1) and amino acid sequence (SEQ ID NO:2) of murine CRP1 (mCRP1). Predicted signal sequence of CRP1 is underlined at the amino-terminus and the experimentally determined pro-peptide cleavage site is indicated by an asterisk. Predicted transmembrane sequence is underlined toward the carboxy-terminus. B) Amino acid alignment of murine CRP1 protein sequence (mCRP1) with murine CD28 (mCD28).

Figure 2. A) DNA (SEQ ID NO:6) and amino acid sequence (SEQ ID NO:7) of murine B7RP1 (mB7RP1). Predicted signal sequence of B7RP1 is underlined at the amino-terminus and the experimentally determined pro-peptide cleavage site is indicated by an asterisk. Predicted transmembrane sequence is underlined toward the carboxy-terminus. B) Amino acid alignment of B7RP1 protein sequence (mB7RP1) with murine CD80 (mCD80).

Figure 3. A) Structure and sequence of the protein coding region of the putative human B7RP1 (hB7RP1; SEQ ID: NO:11 and 12). Predicted signal sequence of hB7RP1 is underlined at the aminoterminus. Predicted signal peptide cleavage sites are marked by asterisks. Predicted transmembrane sequence is underlined toward the carboxy-terminus. B) Amino acid alignment of the putative mature hB7RP1 protein with the mature murine B7RP1 (mB7RP1) protein.

Figure 12. A) Structure and sequence of the protein coding region of human B7RP1 (hB7RP1; SEQ ID NO:16 and 17). Predicted signal sequence of hB7RP1 is underlined at the aminoterminus. Predicted signal peptide cleavage sites are marked by asterisks. Predicted transmembrane sequence is underlined toward the carboxy-terminus. B) Amino acid alignment of the putative mature hB7RP1 protein with the mature murine B7RP1 (mB7RP1) protein.

Figure 13. A) Structure and sequence of the protein coding region of human CRP1 (hCRP1, SEQ ID NO:21 and 22). Predicted signal sequence of hCRP1 is underlined at the amino-terminus. Predicted signal peptide cleavage sites are marked by asterisks. Predicted transmembrane sequence is underlined toward the carboxy-terminus. B) Amino acid alignment of the hCRP1 protein with the murine CRP1 (mB7RP1) protein.